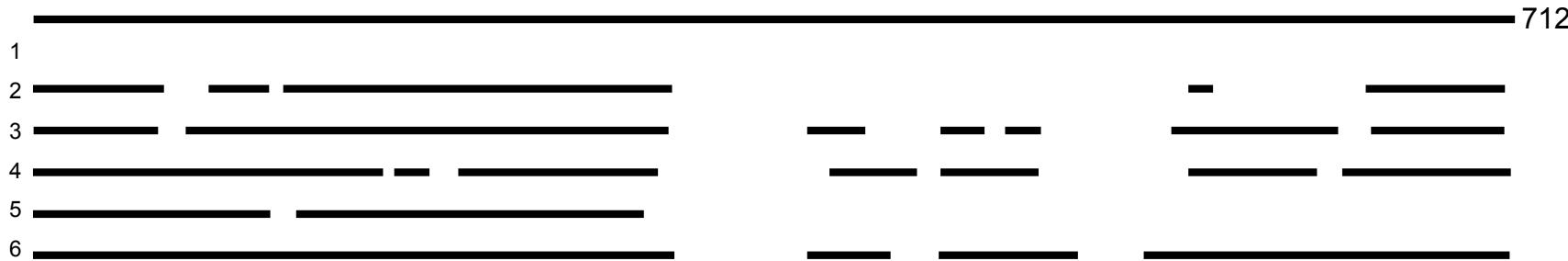
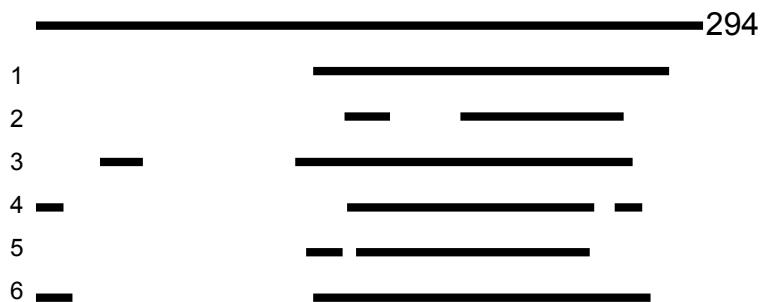


Nucleolin



Nucleophosmin



1. **Ucon** (<http://www.rostlab.org/newwebsite/services/ucon/>)
2. **Disopred2** (<http://bioinf.cs.ucl.ac.uk/disopred/disopred.html>)
3. **RONN** (<http://www.strubi.ox.ac.uk/RONN>)
4. **Disembl: Hot-Loops** (<http://dis.embl.de/>)
5. **Disembl: Remark 465** (<http://dis.embl.de/>)
6. **Disprot: VSL2B** (<http://www.ist.temple.edu/disprot/predictorVSL2.php>)

Supplementary Figure 1. Regions of predicted disorder within nucleolin and nucleophosmin using online resources (1-6). Only regions of predicted disorder ≥ 10 amino acids in length are shown. The protein sequences used for disorder predictions were accession numbers P19338 and P06748 respectively.